IFNFH01	1	AAC								AAT		30
pIFNFH01	1		Met	<u>Thr</u>	Ser	Phe	Asn	Lys	Leu	<u>Asn</u>	Lys	9
IFNFH01	31	CTA	CCA	GGG	ACC	AAC	CCT	GGA	GAA	ACA	GAA	60
plfnfh01	10	Leu	Pro	GJA	Thr	<u>Asn</u>	Pro	Gly	<u>Glu</u>	Thr	<u>Glu</u>	19
IFNFH01	61									TTC		90
pIFNFH01	20	<u> Ile</u>	Cys	<u>Asp</u>	Leu	Leu	Asp	Arg	Glu	<u>Phe</u>	<u>Lys</u>	29
IFNFH01	91									AAA		120
PIENEH01	30	<u>Ile</u>	Ala	<u>Val</u>	<u>Leu</u>	Arg	Lys	Leu	Lys	Lys	Tyr	39
IFNFH01	121									AGA		150
pifnfh01	40	Gln	<u>Asp</u>	Asp	Thr	<u>Glu</u>	Lys	Lys	Phe	Arg	<u>Ile</u>	49
IFNFH01	151									ATT		180
pIFNFH01	50	Leu	Ser	Asp	Lys	Phe	Asn	Lys	Glu	Ile	<u>Glu</u>	59
IFNFH01	181									ATT		210
pIFNFH01	60	Ile	Leu	Lys	Asn	Asn	Gln	Ala	Glu	Ile	Leu	69
IENFH01	211									CTG		240
pIFNFH01	70	<u>Glu</u>	Leu	Lys	Asn	Leu	Thr	Gly	Ile	Leu	Lys	. 79
IFNFH01	241									AGA		270
pIFNFH01	80	Asn	Val	Pro	Gly	Ser	Phe	Asn	Ser	Arg	Ile	89
IFNFH01	271									AAG		300
pifnfh01	90	Asp	GΣΆ	Ala	ГÀЗ	GJA	Arg	Ile	Ser	Lys	Pro	99
IFNFH01	301									CAG		330
pIFNFH01	100	Glu ◀	Asp	Arg	Leu	Phe	Glu	Asn	Thr	Gln	Arg	109
IFNFH01	331	AGA								AAA		360
pIFNFH01	110	Arg	Gln	Lys	Lys	Arg	Asn	Lys	Lys	Lys	stop	118

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IFNFH03	1	AAC	ATG	ACA	TCA	CCA	AAT	GAG	TTA	AAT G	AG	30
pifnfh03	1		<u>Met</u>	<u>Thr</u>	Ser	Pro	Asn	<u>Glu</u>	Leu	Asn (Glu	9
IFNFH03	31	GCA	GCA	GGA	ACT	ACT	CCC	AAA	GAA	ACA G	AG	60
pIFNFH03	10	Ala	Ala	Gly	<u>Thr</u>	Thr	Pro	Lys	Glu	Thr C	<u>Glu</u>	19
IFNFH03	61	ATA	TGT	GAC	ATT	TCA	GAC	AGA	GAA	TTC A	AA	90
pifnfh03	20	<u>Ile</u>	<u>Cys</u>	<u>Asp</u>	Ile	Ser	Asp	Arg	<u>Glu</u>	Phe !	Lys	29
IFNFH03	91	ATA	GCT	TTG	TTG	AAG	AAA	CTC	AAA	GAA A	TT	120
pIFNFH03	30	<u>Ile</u>	Ala	Leu	<u>Leu</u>	<u>Lys</u>	Lys	Leu	Lys	<u>Glu</u>	<u>Ile</u>	39
IFNFH03	121	CAA	GAT	AAT	ACG	GAG	AAG	GAA	CTC	AGA A	TT	150
pIFNFH03	40	<u>Gln</u>	Asp	<u>Asn</u>	Thr	<u>Glu</u>	Lys	Glu	Leu	Arg .	<u>Ile</u>	49
IFNFH03	151	CTA	TCA	GAT	AAA	TTT	AAC	AAG	GAG	ATT C	AA	180
pIFNFH03	50	<u>Leu</u>	<u>Ser</u>	<u>Asp</u>	<u>Lys</u>	Phe	Asn	Lys	Glu	<u>lle</u>	<u>Glu</u>	59
IFNFH03	181	ATG	ATT	AAA	AAG	AAC	CAA	GCA	GAA	ATT (CTG	210
pIFNFH03	60	Met	<u>Ile</u>	Lys	Lys	Asn	Gln	Ala	Glu	Ile	Leu	69
IFNFH03	211	GAG	CTA	AAA	AAT	GCA	GGT	GGC	ATA	TTG A	AAA	240
pIFNFH03	70	<u>Glu</u>	<u>Leu</u>	Lys	<u>Asn</u>	Ala	Gly	Gly	Ile	Leu	Lys	79
IFNFH03	241	ATG	CAT	CAG	AGT	TGG	CTG	GGC	ATG	GTG (3CT	270
pIFNFH03	80	Met	His	Gln	Ser	Trp	Leu	Gly	Met	Val A	Ala	89
IFNFH03	271	CAC	GCC	TGT	AAT	ccc	AGT	ACT	TTG	GGA A	AGC	300
plfnfh03	90	His	Ala	Cys	Asn	Pro	Ser	Thr	Leu	Gly S	Ser	99
IFNFH03	301	CGA	GGT	GGG	TGG	ATC	ACG	AGT	TCA	GGA (GTT	330
pIFNFH03	100	Arg	Gly	Gly	Trp	Ile	Thr	Ser	Ser	Gly /	Val	109
IFNFH03	331	CAA	GAC	CAG	CCT	GGC	CAA	GGC	: AGT	GAA	ACC	360
pIFNFH03	110	Gln	qeA	Gln	Pro	Gly	Gln	Gly	Ser	Glu 3	lhr	119
IFNFH03	361	TCA	TCT	CTA	CTA	AAA	ATA	CAA	AAA	TTA	GCT	390
pIFNFH03	120	Ser	Ser	Leu	Leu	Lys	Ile	Gln	Lys	Leu .	Ala	129
IFNFH03	391		TGC									414
pIFNFH03	130	Gly	Cys	Ser	Gly	Arg	His	Leu	sto	P		136

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IFNFH04	1	AAC	ATG	ACC	TCA	CCA	AAT	GAA	CTA	AAT	AAG	30
pifnfh04	1		<u>Met</u>	Thr	Ser	Pro	Asn	<u>Glu</u>	<u>Leu</u>	<u>Asn</u>	<u>Lys</u>	9
IFNFH04	31	GCA	CCA	GGG	ACC	AAT	CCT	GGA	GAA	ACA	GAG	60
plfnfh04	10	Ala	Pro	Gly	Thr	Asn	Pro	Gly	Glu	Thr	Glu	19
IFNFH04	61	ATG	TAT	GAC	CTT	TCA	GAC	AGA	GAA	TTC	AAA	90
pIFNFH04	20	Met	Tyr	<u>Asp</u>	<u>Leu</u>	Ser	Asp	Arg	<u>Glu</u>	Phe	Lys	29
IFNFH04	91		GCT									120
pIFNFH04	30	Thr	Ala	Ile	Leu	Arg	Lys	Leu	Lys	Glu	<u>lle</u>	39
IFNFH04	121		GAT									150
pIFNFH04	40	<u>Gln</u>	<u>Asp</u>	<u>Asn</u>	Thr	Lys	Lys	Glu	Phe	Arg	Ile	49
IFNFH04	151	CTA	TCA	GAT	AAA	TTT	AAC	AAA	CAG	ATC	GAA	180
pifnfh04	50	Leu	Ser	Asp	Lys	Phe	Asn	Lys	Gln	Ile	Glu	59
IFNFH04	181		ATT									210
pIFNFH04	60	<u>Ile</u>	<u>Ile</u>	Lys	ГЛЗ	Asn	G1n	Ala	Glu	Ile	Leu	69
IFNFH04	211	GAG	CTG	AAA	AAT	GTA	ATT	GAT	ATA	CTA	AAG	240
pIFNFH04	70	<u>Glu</u>	Leu	Lys	Asn	Val	Ile	Asp	Ile	Leu	Lys	79
IFNFH04	241	AAT	GCA	TCA	GTC	TCT	TGA					258
pIFNFH04			Ala									84

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IFNFH08	1	AAC	ATG	ACC	TCA	CCA	AAT	GAA	CTT	AGT	AAG	30
pIFNFH08	1		Met	Thr	Ser	Pro	<u>Asn</u>	<u>Glu</u>	Leu	Ser	Lys	9
IFNFH08	31	GCA	CCA	GGG	ACC	AAT	CAG	GGA	GAA	ACA	GAG	60
pIFNFH08	10	Ala	Pro	Gly	Thr	<u>Asn</u>	Gln	Gly	Glu	Thr	Glu	19
IFNFH08	61	ATA	TAT	GAC	CTT	TCA	GAC	ACA	GAA	TTC	AAA	90
pIFNFH08	20	<u>lle</u>	Tyr	Asp	<u>Leu</u>	Ser	Asp	Thr	Glu	Phe	Lys	29
IFNFH08	91	ATA	GCT	GTT	TTG	AGA	AAC	TCA	AAG	AAG	AAA	120
pIFNFH08	30	<u>Ile</u>	Ala	Val	<u>Leu</u>	Arg	Asn	Ser	Lys	Lys	Lys	39
IFNFH08	121	CTC	AAA	GAA	ATT	CAG	GAT	AAC	ACA	GAG	AAG	150
pIFNFH08	40	<u>Leu</u>	Lys	<u>Glu</u>	<u>Ile</u>	<u>Gln</u>	<u>Asp</u>	Asn	Thr	Glu	<u>Lys</u>	49
IFNFH08	151	GAA	TTC	AGA	ATT	CTA	TCA	GAT	AAA	TTT	AAC	180
pIFNFH08	50	<u>Glu</u>	Phe	<u>Arg</u>	<u>lle</u>	Leu	Ser	Asp	Lys	Phe	neA	59
IFNFH08	181	AAA	GAG	ATT	GAA	ATA	ATT	AAA	AAG	AAT	CAA	210
pIFNFH08	60	Lys	Glu	<u>Ile</u>	<u>Glu</u>	Ile	<u>Ile</u>	Lys	Lys	Asn	Gln	69
IFNFH08	211	GCA	GAA	ATT	CTA	GAG	TTG	AAA	AAT	GCA	ATT	240
pIFNFH08	70	<u>Ala</u>	Glu	Ile	Leu	Glu	Leu	Lys	Asn	Ala	Ile	79
IFNFH08	241	GAC	ATG	CTG	AAT	AAT	GCA	TCA	GAT	TAT	CTT	270
pIFNFH08	80	Asp	Met	Leu	Asn	neA	Ala	Ser	Asp	Tyr	Leu	89
IFNFH08	271				ATT				_			294
pIFNFH08	90	His	Ser	Arg	Ile	Asn	Arg	Asn	sto	p		96

								_				
IENEH10	1	AAC	ATG	ACC	TCA	CCA	AAT	GAG	GTA	AAT .	AAG	30
pIFNFH10	1		Met	Thr	Ser	Pro	<u>Asn</u>	<u>Glu</u>	Val	<u>Asn</u>	<u>Lys</u>	9
											~~~	60
IFNFH10	31									ACG		
pIFNFH10	10	Val	Pro	Met	Thr	<u>Asn</u>	Pro	Gly	Glu	Thr	GIU	19
IFNFH10	61	ΔΤΔ	тст	GAC	СТТ	TCA	GAC	CAA	AAA	TTA	AAA	90
pIFNFH10	20	Tla	Cve	Aen	Len	Ser	Asp	Gln	Lvs	Leu	Lvs	29
breweuro	20	776	Cys	<u> </u>	<u> </u>	<u> </u>		<b>U</b>	-1-			
IFNFH10	91	ATA	GCT	GTG	ATG	AGG	AAA	CTC	AAA	GAA	ATT	120
pIFNFH10	30	Ile	Ala	Val	Met	Arg	Lys	Leu	Lys	Glu	<u>Ile</u>	39
-												
IFNFH10	121									AAA		150
pIFNFH10	40	<u>Gln</u>	Asp	<u>Asn</u>	Thr	Glu	Lys	<u>Glu</u>	Phe	ГÀЗ	<u>Ile</u>	49
			ma3	~~m		mmm	220	222	220	ATT	CCA	180
IFNFH10	151											59
pIFNFH10	50	Leu	Ser	Arg	rya	Pne	ASn	гля	гля	Ile	GLY	33
IFNFH10	181	тта	ATT	GAA	AAT	AAT	CAA	GCA	GAA	ATT	TTG	210
pIFNFH10	60									Ile		69
presento	00	Dea		014								
IFNFH10	211									CTG		240
pIFNFH10	70	Glu	Leu	Lys	Asn	Ala	Ile	<u>Gly</u>	Ile	Leu	Lys	79
-												
IFNFH10	241	AAT	GCA	TCA	GAG	TCC	TTT	AAT	AGC	AAT	ATG	270
pIFNFH10	80	Asn	Ala	Ser	Glu	Ser	Phe	Asn	Ser	Asn	Met	89
									3 CM	C3.C	C/III	300
IFNFH10	271	TAT	CAA	GCA	GAA	GAC	AGA	ATT	AGT	GAG	CTT	99
pifnfh10	90	Tyr	Gln	Ala	Glu	Asp	Arg	He	Ser	Gl u	rea	99
IFNFH10	301	222	ጥአር	»CC	СТЪ	ттт	GAA	ААТ	ACA	CAG	TCA	330
		Tue	THE	Ara	LOU	Phe	Glu	Agn	Thr	Gln	Ser	109
pIFNFH10	100	гÃя	TAT	ALG	Deu	r 114	u					
IFNFH10	331	GAG	GAG	ACC	AAA	AAC	AAC	AAA	AAA	CAA	TGA	360
pIFNFH10	110	Glu	Glu	Thr	Lys	Asn	Asn	Lys	Lys	Gln	stop	118

									<b>•</b>			
IFNFH11	1	CAC	ATG	ACC	TCA	GGA	AAT	GAA	GTA	AAT	AAG	30
pIFNFH11	1									<u>Asn</u>		9
IFNFH11	31	CCA	CCA	GGG	»CC	ልልሞ	<del>ር</del> ጥሞ	GGA	GAA	ACA	GAG	60
	10		Pro									19
pIFNFH11	10	ATG	PLO	GTA	1711	ASII	neu	GIY	910	1111	010	1.7
IFNFH11	61	ATA	TGT	GAC	CTT	TCA	GAT	ACA	GAA	CTC	AGA	90
pIFNFH11	20	<u>lle</u>	Cys	Asp	Leu	Ser	<u>Asp</u>	Thr	<u>Glu</u>	Leu	Arg	29
~~~	91		ACT	CITIC	mm¢.	3.00	***	OTFC	220	CAA	አጥጥ	120
IFNFH11	_				-							39
pIFNFH11	30	<u>116</u>	Thr	vai	Leu	Arg	rĀs	Leu	ASII	Giu	116	39
IFNFH11	121		GAT									150
pIFNFH11	40	Lys	Asp	<u>Asn</u>	<u>Thr</u>	<u>Glu</u>	Met	<u>Glu</u>	Phe	Arg	<u>Ile</u>	49
IFNFH11	151	ጥጥር	TCA	CAT	222	արդ	AAG	222	GAG	ልጥሞ	GAA	180
	50		Ser				_					59
pIFNFH11	50	<u>neu</u>	Ser	Asp	Lys	File	пуэ	nys.	GLU	116	GIU	3,
IFNFH11	181	ATA	ATT	AAA	AGG	AAT	CAA	GCA	GAA	ATT	CTG	210
pIFNFH11	60	<u>Ile</u>	<u> 11e</u>	Lys	Arg	Asn	Gln	Ala	Glu	Ile	Leu	69
IFNFH11	211	GAG	CTG	AAA	AAT	GCA	ATT	GGC	ATA	CTG	AAG	240
pIFNFH11	70	Glu	Leu	Lvs	Asn	Ala	Ile	Glv	Ile	Leu	Lys	79
parmena	, ,											
IFNFH11	241		GCA									270
pIFNFH11	80	Asn	Ala	Ser	Glu	Phe	Leu	Asn	Arg	Arg	Thr	89
IFNFH11	271	GAT	CAA	GCA	GCA	GAA	AAA	TCT	AGT	GAG	CCT	300
pIFNFH11	90		Gln									99
piimimi	30	,,op	02				-10	-				
IFNFH11	301										A GG	330
pIFNFH11	100	Glu	Asp	Arg	Leu	Phe	Glu	Asn	Thr	Gln	Arg	109
IFNFH11	331	ጥርጥ	CAA	AAG	AAA	AAG	ААТ	AAA	AAA	CAA	TAA	360
pIFNFH11	110										stop	118
breugutt	110	Ser	CTII	-133	-15	-73		-10	-10		- JOP	

Figure 7

WO 2004/050702

IFNFH12	1	AAC ATG ACC TCA CCA AAT GAA CTG AAT AAG 3	0
pIFNFH12	1	Met Thr Ser Pro Asn Glu Leu Asn Lys	
IFNFH12	31	CCA CCA GGG ACC AAT CCT GGA GAA ACA GAA 6	_
pIFNFH12	10	Pro Pro Gly Thr Asn Pro Gly Glu Thr Glu 1	9
	61	ATA TGT GAC CTT TCA GAC AAA GAA TTC AAA	0
IFNFH12		AIR 101 GRC 011 101 CIG 121 CIG 121 CIG	9
pIFNFH12	20	Ile Cys Asp Leu Ser Asp Lys Glu Phe Lys 2	9
IFNFH12	91	ATA GCT GTG TTG AAG AAA CTC AAC GAA GCT 12	D
pIFNFH12	30	Ile Ala Val Leu Lys Lys Leu Asn Glu Ala	9
IFNFH12	121	CAA GAT AGC ACA GAG AAG GAA TTC AGA ATT 15	0
pIFNFH12	40	Gln Asp Ser Thr Glu Lys Glu Phe Arg Ile	19
			_
IFNFH12	151	CTA TCA GAT AAA TGT AAC AAA GAC ATT AAA 18	-
pIFNFH12	50	Leu Ser Asp Lys Cys Asn Lys Asp Ile Lys	59
		NUN NUU NAN NAC NAU CAN GCA GAN TUT CTG 21	^
IFNFH12	181	AIA AII AAA AAG AAI CILI COI CILI III CIC	•
pIFNFH12	60	Ile Ile Lys Lys Asn Gln Ala Glu Phe Leu	59
IFNFH12	211	AAG CTG AAA GAT GCA ATT GGA ATA CTG AAG 24	٥
			79
pIFNFH12	70	Lys Leu Lys Asp Ala Ile Gly Ile Leu Lys	_
IFNFH12	241	GAT GCA TCA GAG TTT TTT AAT AGC AGA ACT 2	70
pIFNFH12	80		39
PIENEUIS	60	App Ala bel blu life life inc. box ing life	-
T [7] [7]	271	GAT TGA 27	6
IFNFH12		GAT TOA	-
pIFNFH12	90	Asp stop	9

WO 2004/050702 PCT/EP2003/050939

8/23

				_					_			
IFNFH13	1	AAC		ACC								30
pIFNFH13	1		<u>Met</u>	<u>Thr</u>	<u>Ser</u>	Pro	Asn	<u>Glu</u>	Leu	Asn	<u>Lys</u>	9
IFNFH13	31	GCA	CCA	GGG	ACC	AAT	CCT	GGA	GAA	ACT	GAG	60
pIFNFH13	10	Ala	Pro	Gly	Thr	Asn	Pro	<u>Gly</u>	Glu	Thr	<u>Glu</u>	19
IFNFH13	61			GAC								90
pIFNFH13	20	<u>Ile</u>	Cys	<u>Asp</u>	Leu	<u>Ser</u>	Asp	Arg	r As	Phe	<u>Lys</u>	29
IFNFH13	91	AGA	GCT	GTG	TTG	AAG	AAA	CTC	AAA	GAA	ATT	120
pIFNFH13	30	Arg	Ala	<u>Val</u>	<u>Leu</u>	Lys	ГЛЗ	Leu	Lys	<u>Glu</u>	<u>Ile</u>	39
IFNFH13	121	CAA	AAT	GTÇ	TCA	AAG	AAG	GAA	TTC	AGA	ATT	150
pifnfh13	40	<u>Gln</u>	Asn	Val	Ser	Lys	<u>Lys</u>	<u>Glu</u>	Phe	Arg	<u>Ile</u>	49
IFNFH13	151	CTA	TTA	GAT	AAA	TTT	AAC	AGA	CAG	ATT	GAA	180
pIFNFH13	50	<u>Leu</u>	Leu	Asp	Lys	Phe	Asn	Arg	Gln	Ile	<u>Glu</u>	59
IFNFH13	181	GTA	ATT	AAA	AAT	AAT	CAA	ACA	GAA	ATT	ATG	210
pIFNFH13	60	Val	<u>lle</u>	<u>Lys</u>	Asn	<u>Asn</u>	Gln	Thr	<u>Glu</u>	Ile	Met	69
IFNEH13	211			AAA								240
pIFNFH13	70	<u>Glu</u>	Leu	Lys	<u>Asn</u>	<u>Ala</u>	Ile	<u>Gly</u>	Ile	Leu	Lys	79
IFNFH13	241			CAG								270
pIFNFH13	80	Met	His	Gln	Ser	Ser	Leu	Ile	Ala	Ala	Leu	89
IFNFH13	271	ATC	AAA	CAG	AAG	AAA	G AA	TTA	GTG	AAC	CTG	300
pieneh13	90	Ile	Lys	Gln	ГÀ2	ГÀЗ	Glu	Leu	Val	Asn	Leu	99
IFNFH13	301			GCC								330
pIFNFH13	100	Lys	Thr	Ala	Tyr	Leu	Lys	Ile	His	Arg	Gly	109
IFNFH13	331	GAC	AAA	AGA	AAA	AAA	TAT	AAA	AGA	ATG	AA G	360
pIFNFH13	110	•	Lys	Arg	Lys	Lys	Tyr	Lys	Arg	Met	Lys	119
IFNFH13	361	CAC	ACC	TAA	•							369
pIFNFH13				stop	•	,						121

IFNFH14 pIFNFH14	1	AAC	ATG Met	ACA <u>Thr</u>	TCA Ser	ACA Thr	AAG Lys	GAA Glu	CT A Leu	AAT Asn	AAG Lys	30 9
IFNFH14 pIFNFH14	31 10					AAT Asn						60 19
IFNFH14 pIFNFH14	61 20					TTA Leu						90 29
IFNFH14 pIFNFH14	91 30					AGG Arg						120 39
IFNFH14 pIFNFH14	121 40					GAG Glu						150 49
IFNFH14 PIFNFH14	151 50	CTA Leu	TCA Ser	GAT Asp	AAA Lys	TTT Phe	AAC Asn	AAA Lys	GAG Glu	Ile	GAA Glu	180 59
IFNFH14 pIFNFH14	181 60	ATA Ile	ATT Ile	AAA Lys	AAG Lys	AAT Asn	CAA Gln	GCA Ala	GAA Glu	ACT Thr	CTG Leu	210 69
IFNFH14 pIFNFH14	211 70	GAG Glu	CTA Leu	AAA Lys	TAA neA	GCA Ala	GTT Val	GGC G1y	ACA Thr	CTA Leu	ACA Thr	240 79
IFNFH14 pIFNFH14	241 80					TCC Ser						270 89
IFNFH14 pIFNFH14	271 90	GAT Asp	ATA Ile	GCA Ala	GAA Glu	AGA Arg	AGA Arg	ATT Ile	AGT Ser	GAA Glu	CTT Leu	300 99
IFNFH14 pIFNFH14		AAA Lys	GAC Asp	AGG Arg	CTA Leu	TTT Phe	GAA Glu	AAT Asn	ACA Thr	GTC Val	AGA Arg	330 109
IFNFH14 pIFNFH14						ATA						348 114

IFNFH15	1	AAT	ATG									30
pIFNFH15	1		<u>Met</u>	<u>Thr</u>	Ser	Pro	<u>Asn</u>	<u>Glu</u>	Leu	Asn	Lys	9
IFNFH15	31	GCA	CCA	GGG	ATC	AAT	CCT	GGG	GAA	ACA	GAA	60
pIFNFH15	10	Ala	Pro	Gly	Ile	<u>Asn</u>	Pro	<u>Gly</u>	<u>Glu</u>	<u>Thr</u>	<u>Glu</u>	19
IFNFH15	61		TGT									90
pIFNFH15	20	Ile	Cys	<u>Asp</u>	<u>Leu</u>	Ser	Asp	Arg	Glu	Phe	Thr	29
IFNFH15	91	ATA	GCT	GTT	TCG	AGG	AAG	CTA	AAC	AAA	ATC	120
pIFNFH15	30	<u>Ile</u>	Ala	<u>Val</u>	Ser	Arg	Lys	Leu	Asn	ГÀЗ	<u>lle</u>	39
IFNFH15	121	CAA	GAT	AAC	ATG	GAG	AAG	GAA	TTC	AGA	ATC	150
pIFNFH15	40	<u>Gln</u>	Asp	Asn	Met	<u>Glu</u>	Lys	<u>Glu</u>	Phe	Arg	<u>Ile</u>	49
IFNFH15	151	CTA	TCA	GAT	AAA	TTT	AAT	GAA	GAG	ATT	GAA	180
pIFNFH15		Leu	Ser	Asp	Lys	Phe	Asn	Glu	Glu	<u>Ile</u>	<u>Glu</u>	59
IFNFH15	.181	ATA	ATT	AAA	AAG	AAT	CAA	GCA	GAA	ATT	CTG	210
pIFNFH15		Ile	<u>Ile</u>	<u>Lys</u>	Lys	<u>Aşn</u>	Gln	<u>Ala</u>	<u>Glu</u>	<u>Ile</u>	Leu	69
IFNFH15	211	GAG	CTG	AAA	AAC	GCA	ATT	GAC	ATG	TTG	AAG	2 40
pIFNFH15			<u>Leu</u>									79
IFNFH15	241	AAT	GCA	TCA	GAG	AAT	CTC	ACC	AGC	AGA	ACT	270
pIFNFH15			Ala									89
IFNFH15	271	GAT	CAA	GCA	AGA	GAA	ATA	ATT	AGT	AAG	CTT	300
pIFNFH15		Asp	Gln	Ala	Arg	Glu	Ile	Ile	Ser	Lys	Leu	99
IFNFH15	301	GAA	GAC	AGG	CTA	TTT	GAA	AAC	ACA	AAG	TCA	330
pIFNFH15	100	Glu	Asp	Arg	Leu	Phe	Glu	Asn	Thr	Lys	Ser	109
IFNFH15	331	GAG	GAG	ACA	AAT	GGA	AAA	AGA	ATA	AAA	TGC	360
pifnfh15		Glu	Glu	Thr	neA	Gly	Lys	Arg	Ile	Lys	Cys	119
IFNFH15	361	AAT	GAA	GCA	CAC	CTA	CAA	GAA	CTA	GAA	AAT	390
pIFNFH15	120	Asn	Glu	Ala	His	Leu	Gln	Glu	Leu	Glu	Asn	129
IFNFH15	391		TTC									420
pIFNFH15			Phe									139
IFNFH15	421		CTT	AAA	TAG							432
pIFNFH15	140	Gly	Leu	Lys	sto	p						142

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IFNFH20	1	AAC	ATG	CCC	TTA	CCA	AAT	GAG	CTA	AAT	AAG	30
pIFNFH20	1		<u>Met</u>	Pro	Leu	Pro	<u>Asn</u>	<u>Glu</u>	<u>Leu</u>	Asn	<u>Lys</u>	9
IFNFH20	31	GCG	CCA	GGG	ACC	AAT	CCT	GGA	GAA	ACA	GAG	60
pIFNFH20	10	Ala	Pro	Gly	Thr	Asn	Pro	<u>Gly</u>	Glu	Thr	Glu	19
IFNFH20	61				CTT							90
pIFNFH20	20	Thr	Сув	Asp	<u>Leu</u>	<u>Ser</u>	Asp	Arg	Glu	Phe	Lys	29
IFNFH20	91	ATA	GCT	GTG	TTG	AGA	AAA	CTC	AAA	GAA	ATT	120
pIFNFH20	30	<u>Ile</u>	Ala	<u>Val</u>	Leu	Arg	Lys	Leu	Lys	<u>Glu</u>	<u>Ile</u>	39
IFNFH20	121	CAA	GAG	AAT	ACA	GAC	AAG	GAA	TTG	AGA	ATT	150
pIFNFH20	40	Gln	Glu	<u>Asn</u>	Thr	Asp	Lys	<u>Glu</u>	Leu	Arg	Ile	49
IFNFH20	151	CTA	TCA	GAT	AAA	TTT	AAC	AAA	GAA	ATC	AAA	180
pIFNFH20	50	Leu	Ser	Asp	<u>Lys</u>	Phe	Asn	Lys	Glu	Ile	Lys	59
IFNFH20	181	ATA	ATG	AAA	AAG	AAT	CAA	GCA	GAA	ATT	CTG	210
pIFNFH20	60	Ile	Met	Lys	Lys	Asn	Gln	Ala	Glu	<u>Ile</u>	Leu	69
IFNFH20	211	AAG	CTG	AAA	AAT	TCA	ATT	AGT	ATA	ATG	AAG	240
pIFNFH20	70	Lys	Leu	Lys	Asn	Ser	Ile	Ser	Ile	Met	Lys	79
IFNFH20	241	AAT	GCA	TCA	TAG							252
pIFNFH20	80	Asn	Ala	Ser	stop	9						82

IFNFH23	1	AAC	ATG	ACC	TCA	CCA	AAT	GAA	CTG	AAT	AAG	30
pIFNFH23	1				Ser							9
•								_				
IFNFH23	31	GCA	CCA	GGG	ACG	AAT	TTA	GGA	GAA	ACA	GAG	60
pIFNFH23	10	Ala	Pro	Glv	Thr	Asn	Leu	Gly	Glu	Thr	Glu	19
•				_								
IFNFH23	61	ATT	TGT	GAC	CTT	TCA	GAC	AGA	GAA	TTC	AAG	90
pIFNFH23	20	Ile	Cvs	Asp	Leu	Ser	Asp	Arg	Glu	Phe	Lys	29
•								•				
IFNFH23	91	AAA	GCT	GTG	TTG	CAG	AAG	CTC	AAA	GAA	ATT	120
PIFNFH23	30	Lys	Ala	Val	<u>Leu</u>	Gln	Lys	Leu	Lys	Glu	Ile	39
		_					_					
IFNFH23	121	CAA	GAT	AAC	ACA	GAG	AAG	GAG	TTC	AGA	ATT	150
pIFNFH23	40	Gln	Asp	Asn	Thr	<u>Glu</u>	Lys	Glu	Phe	Arq	<u> Ile</u>	49
IFNFH23	151	CTA	TTA	CAT	AAA	TTT	AAC	AAA	GAG	ATT	AAA	180
pIFNFH23	50	Leu	Leu	His	Lys	Phe	Asn	Lys	Glu	Ile	Lys	59
IFNFH23	181				AAG							210
pIFNFH23	60	Ile	Ile	Lys	Lys	<u>Asn</u>	<u>Gl.n</u>	Ala	Glu	Ile	Leu	69
IFNFH23	211	-			AAT		-			_	_	240
pIFNFH23	70	<u>Glu</u>	Ala	Lys	<u>Asn</u>	<u>Ala</u>	Thr	Asp	Ile	Leu	Met	79
IFNFH23	241				GAC							270
PIFNFH	80	Asn	Ala	Ser	Asp	Pro	Ile	A sn	Ser	Thr	Ile	89
IFNFH23	271				GAA				-		_	300
PIENEH	90	Asp	Glu	Ala	Glu	Glu	Arg	IJe	Ser	Glu	Leu	99
			4	_								
IFNFH23	301				CTA							327
pIFNFH23	100	Glu	Asp	Arg	Leu	Phe	Glu	Ser	Ile	stop	•	1 07

IFNFH25 pIFNFH25	1	AAC			CCA Pro				AAG Lys	30 9
IFNFH25 pIFNFH25	31 10			 	AAT <u>Asn</u>				GAG Glu	60 19
IFNFH25 pIFNFH25	61 20			 	TCA <u>Ser</u>		_		AAA Lys	90 29
IFNFH25 pIFNFH25	91 30				AGG Arg					120 39
IFNFH25 pIFNFH25	121 40				GAG <u>Glu</u>				Ile	150 49
IFNFH25 pIFNFH25	151 50				TTT Phe				GAA Glu	180 59
IFNFH25 pIFNFH25	181 60				AAT <u>Asn</u>				CCG Pro	210 69
IFNFH25 pIFNFH25	211 70				GCA Ala					240 79
IFNFH25 pIFNFH25	241 80			 	TCT Ser			 	ATT Ile	270 89
IFNFH25 pIFNFH25	271 90		CAA Gln		TAA stop	,				285 93

WO 2004/050702 PCT/EP2003/050939

14/23

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IFNFH27	1	AAC	ATG	ACC	TCG	CCT	AAT	GAA	CTA	AAT	GAA	30
pIFNFH27	1		<u>Met</u>	Thr	<u>Ser</u>	Pro	<u>Asn</u>	<u>Glu</u>	Leu	Asn	Glu	9
IFNFH27	31		CCA									60
pIFNFH27	10	Ala	Pro	Gly	Thr	Asn	Pro	Ala	<u>Glu</u>	Thr	<u>Glu</u>	19
IFNFH27	61	ATA	TGT	AAC	ATT	TTA	GAC	AGA	GAA	TTC	AAA	90
pIFNFH27	20	<u>Ile</u>	<u>Cys</u>	Asn	Ile	Leu	Asp	Arg	Glu	Phe	Lys	29
IFNFH27	91		GCT									120
pIFNFH27	30	<u>Ile</u>	Ala	<u>Val</u>	Leu	Arg	Lys	Leu	Asn	<u>Glu</u>	Ile	39
IFNFH27	121		GAT									150
pIFNFH27	40		<u>Asp</u>		_							49
IFNFH27	151		TCA									180
pIFNFH27	50		Ser		_							59
IFNFH27	181		ATT				-					210
pIFNFH27	60		<u>Ile</u>									69
IFNFH27	211		TTG									240
pIFNFH27	70	<u>Glu</u>	Leu	Lys	Asn	Ala	Thr	Asp	Ile	Arg	Lys	79
IFNFH27			GCA									270
pIFNFH27	80	Asn	Ala	Ser	Gly	Ser	Leu	Asn	Lys	Ar g	Leu	89
IFNFH27	271		CTT									300
pIFNFH27	90	neA	Leu	Ser	Glu	Glu	Arg	Ile	Ser	Glu ◀	Leu	99
IFNFH27	301		GAT									330
pIFNFH27	100	<u></u>	Asp				Asp	Asn	Ile	Gln	Ser	109
IFNFH27	331		GAA									345
pIFNFH27	110	Glu	Glu	Ala	Asn	sto	P					113

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IFNFH31	1	AAT	ATG	ACC	TCA	CCA	AAT	GAA	CTA	AAT	AAG	30
pIFNFH31	1		<u>Met</u>	Thr	Ser	Pro	<u>Asn</u>	<u>Glu</u>	Leu	Asn	Lys	9
IFNFH31	31	GTA	CCA	GGG	GCC	AAT	CCT	GGA	GAA	ACA	GAG	60
pifnfh31	10	Val	Pro	GJ À	Ala	Asn	Pro	<u>Gly</u>	<u>Glu</u>	Thr	<u>Glu</u>	19
IFNFH31	61		TGT									90
pIFNFH31	20	<u>Ile</u>	CAa	Asp	His	Ser	Glu	Arg	<u>Glu</u>	Phe	Lys	29
IFNFH31	91	ATA	ACT	GTC	TTG	AGG	AAA	CTC	AAA	GAC	ATT	120
pIFNFH31	30	<u>Ile</u>	Thr	<u>Val</u>	Leu	Arg	Lys	Leu	Lys	Asp	Ile	39
IFNFH31	121		GAT									150
pIFNFH31	40	His	<u>Asp</u>	<u>Asn</u>	Thr	<u>Glu</u>	Lys	Thr	Ile	Arg	<u>Ile</u>	49
IFNFH31	151		TCA									180
pifnfh31	50	Leu	Ser	Asp	Lys	Phe	Asn	Lys	Asp	Ile	Glu	59
IFNFH31	181	ATA	ATT	TTA	AAA	AAT	CAA	GAT	GAT	ATT	CTG	210
pIFNFH31	60	<u>Ile</u>	<u>Ile</u>	Leu	Lys	Asn	<u>Gln</u>	Asp	Asp	Ile	Leu	69
IFNFH31	211		CTG						_			240
pIFNFH31	70		Leu									79
IFNFH31	241		GAA		_							270
pIFNFH31	80		Glu		•					-		89
	271		GAA									300
pIFNFH31	90	_	Glu						_	-		99
IFNFH31	301		GAC									330
pIFNFH31	100	Glu	Asp	Ser	Leu	Phe	Glu	Asn	Ile	Gln	Ser ∢	109
IFNFH31	331		AAG									360
pIFNFH31	110	Glu	Lys	Lys	Ala	Lys	Lys	Val	Lys	Gln	Thr	119
IFNFH31			AAA									384
pIFNFH31	120	Asn	Lys	Lys	Arg	Ser	Met	Tyr	sto	•		126

IFNFH32	1	AAC								AAA A		30
pifnfh32	1		Met	Thr	Ser	Pro	Asn	Lys	Leu	Lys <u>I</u>	ys	9
IFNFH32	31	GCA	CCA	GGG	ACC	AAT	CCT	GGA	GAA	ACA G	AA	60
pIFNFH32	10	Ala	Pro	Gly	<u>Thr</u>	<u>Asn</u>	Pro	<u>Gly</u>	<u>Glu</u>	Thr G	<u>Slu</u>	19
IFNFH32	61	ACA	тст	GGA	CTT	TCA	CAG	AGA	GAA	TTC A	AA	90
pIFNFH32	20									Phe I		29
********	01	CIDA	CCTT	CTC	marc:	N.C.C.	***	СТС	444	GAA A	тт	120
IFNFH32	91 30									Glu 1		39
pIFNFH32	30	AGT	ATG	<u>va1</u>	Tie G	ALG	Dy 5	<u> </u>	-10	<u> </u>		
IFNFH32	121									AGA A		150
pIFNFH32	40	<u>Gln</u>	Asp	<u>Asn</u>	Arg	<u>Glu</u>	Lys	Glu	Phe	Arg 1	<u>lle</u>	49
IFNFH32	151	CTA	TCA	СРТ	AAA	ттт	AAC	AAA	GAG	ATT G	AA	180
pIFNFH32	50	Val	Ser	Asp	Lvs	Phe	Asn	Lvs	Glu	Ile C	Glu	59
printinge	50		<u> </u>		=1.				•			
IFNFH32	181	ATA	ATT	AAA	AAG	AAT	CAG	GCA	GAA	ATA C	TG	210
pIFNFH32	60	Ile	<u>Ile</u>	Lys	Lys	Asn	Gln	<u>Ala</u>	Glu	Ile 1	Leu	69
_		_	<u> </u>									
IFNFH32	211		CTG									237
pIFNFH32	70	<u>Gl.u</u>	Leu	Lys	Asn	Gln	Leu	Ala	Tyr	stop		77

IFNFH36	1	AAC	ATG	ACC	TCA	CCA	AAC	AAA	CTA	AAT	AAG	30
pifnfh36	1		<u>Met</u>	<u>Thr</u>	Ser	<u>Pro</u>	<u>Asn</u>	Lys	Leu	<u>Asn</u>	<u>Lys</u>	9
IFNFH36	31	GCA	CCC	AGG	GCC	AAT	TCT	GGA	GAA	ACA	GAG	60
pifnfh36	10	Ala	Pro	Arg	Ala	<u>Asn</u>	Ser	<u>Gly</u>	<u>Glu</u>	Thr	<u>Glu</u>	19
IFNFH36	61	ATA	CGT	AAA	CTT	TCA	AAC	ACA	GAA	ATC	AAG	90
pifneh36	20	<u>Ile</u>	Arg	Lys	Leu	<u>Ser</u>	Asn	The	<u>Glu</u>	Ile	<u>Lys</u>	29
IFNFH36	91	ATA	GCT	GTG	TTG	AGA	AAA	CTC	AAA	GAA	ATT	120
pifnfh36	30	<u>Ile</u>	Ala	<u>Val</u>	Leu	Arg	Lys	Leu	Lys	Glu	<u>Ile</u>	39
IFNFH36	121	CAA	GAT	AAC	ACA	GAG	AAA	GAA	TTC	AGA	ATT	150
pifnfh36	40	<u>Gln</u>	<u>Asp</u>	<u>Asn</u>	Thr	<u>Glu</u>	Lys	<u>Glu</u>	Phe	Arg	<u> Ile</u>	49
IFNFH36	151									ATT		180
pIFNFH36	50	Leu	Ser	<u>Asp</u>	Lys	Phe	Asn	Lys	Glu	Ile	<u>Glu</u>	59
IFNFH36	181	ATA	ACT	AAA	AAG	AAT	CAA	GCA	GAA	ATT	CTG	210
plfnfh36	· 60	<u> Ile</u>	Thr	Lys	Lys	<u>Asn</u>	<u>Gln</u>	Ala	Glu	Ile	Leu	69
IFNFH36	211									CTG		240
pIFNFH36	70	<u>Glu</u>	<u>Leu</u>	Arg	Asn	<u>Ala</u>	Ile	Asp	Ile	Leu	Lys	79
IFNFH36	241									AGA		270
plfnfh36	80	Asn	Ala	Ser	GЉ	Ser	Phe	Asn	Ser	Arg	Ile	89
IFNFH36	271	GAG	CAA	GCA	GAA	TAA	•					285
pifnfh36	90	Glu	Gln	Ala	Glu	stor	>					93

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IFNFH37	1	AAC	ATG	ACC	TCA	CTA	AAT	GAA	CTA	AAT	AAG	30
pIFNFH37	1										Lys	9
IFNFH37	31	GCA	CCA	GGG	GCC	AAC	CCT	GGA	GAA	ACA	GAG	60
pIFNFH37	10	Ala	Pro	Gly	Ala	Asn	Pro	Gly	Glu	Thr	Glu	19
IFNFH37	61	ATA	TGC	GAC	CTT	TCA	GAC	AGA	GAA	TTC	AAA	90
PIFNFH37	20	<u>Ile</u>	<u>Cys</u>	Asp	Leu	Ser	Asp	Arg	<u>Glu</u>	Phe	Lys	29
IFNFH37	91	ATA	GCT	GTG	TTG	GGG	AAA	TTC	AAA	GAT	AAC	120
oIFNFH37	30	Ile	Ala	<u>Val</u>	Leu	Gly	Lys	Phe	Lys	Asp	Asn	39
FNFH37	121	ACA	GAG	AAG	GAA	TTC	AGA	ATT	CTA	TCA	GAT	150
oIFNFH37	40	<u>Thr</u>	<u>Glu</u>	Lys	<u>Glu</u>	Phe	Arg	<u>lle</u>	Leu	Ser	Asp	49
FNFH37	151	AAA	TTT	AAC	AAA	GAG	ATT	GAA	ATA	ATT	AAA	180
oIFNFH37	50	<u>Lys</u>	Phe	Asn	Lys	Glu	<u>Ile</u>	<u>Glu</u>	<u>Ile</u>	Ile	Lys	59
FNFH37	181	AAG	AAT	CAA	GCA	GAA	ATT	CTG	GAG	CTG	AAA	210
oIFNFH37	60					<u>Glu</u>						69
FNFH37	211	AAT	GCA	ATT	GCC	ACA	TTA	AAG	AAT	GCA	TTA	240
oIFNFH37	70	Asn	<u>Ala</u>	Ile	Ala	Thr	Leu	Lys	Asn	Ala	Leu	79
FNFH37	241	GAG	TTT	TTT	AAT	AGC	AGA	ATT	TAT	GGA	GCA	270
IFNFH37	80	Glu	Phe	Phe	Asn	Ser	Arg	Ile	Tyr	Gly	Ala	89
FNFH37	271	GĀA	AAA	AAG	AAT	TAG						285
IFNFH37	90	Glu	Lys	Lys	Asn	stop)					93

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IFNFH39	1	TCA	ATG	GCC	AGA	CAC	CTA	CAA	ACA	TCC	ACT	30
pIFNFH39	1					His	-					9
•	_				5			U				•
IFNFH39	31	AGC	ATC	AAG	ACC	ATC	CAG	GAA	AAT	AGG	ACC	60
pIFNFH39	10	Ser	Ile	Lys	Thr	Ile	Gln	Glu	Asn	Arq	Thr	19
-				-				_		•		
IFNFH39	61	TCA	CCA	AGT	GAA	CTA	AAT	AAG	GCA	CCA	GGG	90
pIFNFH39	20	Ser	Pro	Ser	Glu	Leu	Asn	Lys	Ala	Pro	Gly	29
			_								-	
IFNFH39	91	GCC	AGT	CTT	GGA	GAA	ACA	GAG	ATA	TGT	GAT	120
pifnfh39	30	Ala	Ser	Leu	Gly	<u>Glu</u>	Thr	<u>Glu</u>	Ile	Cys	Asp	39
IFNFH39			-			GAA						150
pIFNFH39	40	Leu	Ser	Asn	Arg	<u>Glu</u>	Leu	Lys	Ile	Ala	<u>Val</u>	49
IFNFH39						AAA				-		180
pIFNFH39	50	<u>Leu</u>	Arg	Lys	Leu	Lys	<u>Glu</u>	<u>Ile</u>	Gln	Asp	Ser	59
IFNFH39	181					TTC						210
pIFNFH39	60	Thr	Glu	<u>Lys</u>	Glu	Phe	Arg	Ile	Leu	Ser	Asp	69
IFNFH39	211		mmm									0.40
pIFNFH39	70					CAA						240
brene H23	70	rAa	Pne	ASI	rAa	<u>Gln</u>	11e	GIU	TTG	116	ràs	79
IFNFH39	241	770	N C TP	CAA	CCA	GAA	א חימי	CEC	CAC	OFF		270
pIFNFH39						Glu						89
PILMENSS	00	ASII	Ser	Gilli	WIG	GIU	116	Ten	Gita	neu	Lys	09
IFNFH39	271	ААТ	GCA	АТТ	GAC	TTA	CTG	AAG	аат	CCA	тсъ	300
pIFNFH39						Leu						99
F								~,,				,,,
IFNFH39	301	GAA	TCT	CCT	AAT	AGT	AGA	ATT	AAT	CAA	GTA	330
pIFNFH39	100	Glu	Ser	Pro	Asn	Ser	Arq	Ile	Asn	Gln	Val	109
-		4-					•	_				
IFNFH39	331	GAA	GAA	TGA								339
pIFNFH39	110	Glu	Glu	stop	,							111
				_								

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IFNFH42	1	TCA	ATG	CCA	AGA	CAC	CAA	AGA	ACA	CCT	ACT	30
pIFNFH42	1		Met	Pro	Arg	His	Gln	Arg	Thr	Pro	Thr	9
-								_				
IFNFH42	31		ATC	_								60
pifnfh42	10	Arg	Ile	Asn	Thr	Ile	Gln	Glu	Asn	Thr	Thr	19
												••
IFNFH42	61		TCA									90
pIFNFH42	20	Ser	ser	Asn	GIU	Leu	Asn	GIU	Ата	Pro	Gly	29
IFNFH42	91	አጥር	ACT	ССТ	CCA	CAA	7 C7	CAG	እጥአ	TCT.	CAC	120
pIFNFH42	30										Asp	39
PILMENAL	50	*16	TIIL	110	917	Giu	1111	914	110	د بري	<u>vash</u>	3,
IFNFH42	121	CTT	TCA	GAC	AGA	GAA	TTC	AAA	GTA	GCT	GTG	150
pIFNFH42	40	Leu	Ser	Asp	Arq	Glu	Phe	Lvs	Val	Ala	Val	49
•					-				•			
IFNFH42	151	TTG	AGA	GAG	CTC	AAA	GAA	ATT	CAA	GAT	AAC	180
pIFNFH42	50	Leu	Arg	Glu	Leu	Lys	Glu	Ile	<u>Gln</u>	Asp	Asn	59
IFNFH42	181		GAG									210
pIFNFH42	60	<u>Thr</u>	<u>Glu</u>	Lys	Lys	Phe	Arg	Ile	Leu	Pro	<u>Asp</u>	69
												0.40
IFNFH42			TTT									240
pIFNFH42	70	rys	Pne	тте	гĀЗ	GLU	TIE	GIU	11e	Tre	Lys	79
IFNFH42	241	AAC	ААТ	CAA	ምር አ	CAA	מייד מ	CTC	CNC	CTC	222	270
pIFNFH42	80										Lys	89
prenenaz	00	цуз	ASII	9111	Ser	910	116	Беи	910	<u>neu</u>	DAS	0,5
IFNFH42	271	AAC	CCA	ACT	GCT	GTA	CTG	AAG	AAT	GCA	TCA	300
pIFNFH42			Pro		_	-						99
•												
IFNFH42	301	GAG	TCC	CTT	AAT	AGC	AGA	ATG	GAT	CGA	GTA	330
pIFNFH42	100	Glu	Ser	Leu	Asn	Ser	Arg	Met	Asp	Arg	Val	109
		←										
IFNFH42	331		AAG									345
pIFNFH42	110	Glu	Lys	Lys	Asn	sto	.					113

PCT/EP2003/050939

21/23

WO 2004/050702

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atsphelnk lpwtnpgeteicdlsd tefkisvlk<u>n</u>----lkei
INSP037
                                             TSPNELNK APGTNPGETEICDLSD REFRIAVLRK---- LKEI
PIFNFHcon
                                             MTSPNELNKAPG_INPGETEICDLSDREF<u>T</u>IAV<u>S</u>RK----L<u>NK</u>I
MTSPN_KL<u>K</u>KAPGTNPGETE<u>T</u>CGLSQREFKVAVLRK----LKEI
pIFNFH15
pIFNFH32
                                             ats lnelnkapganpgeteïcdlsdrefkiavlgk----
pIFNFH37
                                             HTSPNELNKAPGTNPGET EMYDLSDREFK TAILRK---- LKEI
pIFNFH04
                                             MTSPNELN <u>EAAGTTPK</u>ETEICDISDREFKIALLKK ---- LKEI
PIFNFH03
                                             MTSPNEL <u>Skapgtnogetei y</u>dlsd<u>tefkiavlr</u>nskkklkei
M <u>pl</u>pnelnkapgtnogete <u>t</u>cdlsdrefkiavlrk----lkei
pifnfh08
PIFNFH20
DIFNFH23
                                             MTSPNELNKAPGTN LGETEICDLSDREFK KAVLQK----LKEI
                                             ntspnelnk <u>p</u>pgtn<del>p</del>geteicdlsdkefkiavlkk ---- l<u>n</u>ea
pIFNFH12
                                             n <u>aspnk</u>lnkap<u>e</u>tnp<u>k</u>etevcdlsdre <u>l</u>ki<u>p</u>vlrk----<u>fn</u>ei
pIFNFH25
                                             NTSPNELNKAPGTNPGETEICDLSDR KFKRAVLKK---- LKEI
pIFNFH13
                                             ts <u>tkelnkapvn</u>npgetelcdl<u>l</u>dk<u>k</u>fkiavlrk----lk<u>g</u>i
pIFNFH14
                                              TSPN KLNKAPRANSGETEIRKLSNTEIKIAVLRK----LKEI
pIFNFH36
pifnfh39
                 MARHLQTSTSIKTIQEN
                                             RTSPSELNKAPGASLGETEICDLS NRELKIAVLRK---- LKEI
                                             MTS GNEVNKAPGTN LGETEICDLSD TELRITVLRK---- LNEI
pIFNFH11
                                             NTSPNELN EAPGTNPAETEICNILDREFKIÄVLRK----LNEI
pIFNFH27
                 MTSPN KLNKLPGTNPGETEICDL LDREFKIAVLRK ---- LKKY
MTSPNELNK VPGANPGETEICDL SCREFKIAVLRK ---- LKDI
MTSPNEVNK VPMTNPGETEICDLSD QKLKIAVMRK ---- LKEI
MPRHQRTPTRINTIQEN TTSSNELNEAPGITPGETEICDLSDREFKVAVLR E---- LKEI
pIFNFH01
pIFNFH31
pIFNFH10
pIFNFH42
INSP037
                 Odnieke Skirsdka kkoje i ikonove i petuvy d
pIFNFHcon
                 QDNTEKE PRILSDKFNKEIEIIKKNQAEILELKNAI
                 QDN MEKEFRILSDKFN EEIEIIKKNQAEILELKNAI MLKNASENLTSRTDQAREIISKL
pIFNFH15
                 QDN REKEFRIVSDKFNKEIEIIKKNQAEILELKN QUAY
pIFNFH32
                  KONTEKEFRILSDKFNKEIEIIKKNQAEILELKNAIATLKNALEFFNSRIYGAEKKN
pIFNFH37
                  QDNT KKEFRILSDKFNK QIEIIKKNQAEILELKN VIDILKNASVS
pIFNFH04
                 ODNTEKE LRILSDKFNKËTEMIKKNOAETLELKNA GGILKMHOSWLGMVAHACNPSTLGS
ODNTEKEFRILSDKFNKETETIKKNOAETLELKNATUMLNNASDYLHSRINKN
pIFNFH03
PIENEH08
                  QENTOKE LRILSOKFNKEI KIMKKNQAEIL KLKNSISIMKNAS
pIFNFH20
                 QDNTEKEFRIL LHKFNKEI KIIKKNQAEILE AKNATDILMNASDPINSTID EAEERISEL
QD STEKEFRILSDK CNKDI KIIKKNQAE FLKLKDAIGILKDASEFFNSRTD
QDNTEKEFRILSDKFNKEIEIIKKNQAEI PEVKNAINTLKNSSESLNSRIDQAE
QNVSKKEFRILLDKFNRQIEVIKNNQTEIMELKNAIGILKMHQSSLIAALIKQKKELVNL
pIFNFH23
pIFNFH12
pIFNFH25
pIFNFH13
                 Q NVSKKEFRILLDKFNRQIEVIKNNQTEIMELKNAIGILKMHQSSLIAALIKQKKELVNL
Q NNTEKEFRILSDKFNKEIEIIKKNQAE TLELKNAVGTLTKASQSFKSRMDIAERRISEL
QDNTEKEFRILSDKFNKEIEI TKKNQAEILELKNAIDILKNASGSFNSRIEQAE
QD STEKEFRILSDKFNK QIEIIKNSQAEILELKNAIDLLKNASESPNSRINQVEE
KDNTEMEFRILSDKF KKEIEIIKRNQAEILELKNAIGILKNASESPNSRINQVEE
QDDTEKE LKVLSDKIIKEIEIIKMQAEILELKNA TDIRKNASGSLNKRLNLSEERISEL
QD DTEKKFRILSDKFNKEIEIIK NNQAEILELKNA TDIRKNASGSLNKRLNLSEERISEL
HDNTEKTIRILSDKFNKDIEII LKNQDDILELENAIGULKNESGFFNSRMDEAEEIIRKL
ODNTEKEFKILS RKFNKKIGLIENNOAEILELKNAIGILKNASSSENSN MYOAEDRISEL
pIFNFH14
pIFNFH36
pIFNFH39
pIFNFH11
pifnfh27
pIFNFH01
pIFNFH31
                 QDNTEKEFKILS RKFNKKIGLIENNQAEILELKNAISILKNASESFNSN MYQAEDRISEL
QDNTEK KFRILPDKFIKEIEIIKKNQSEILELKN PTAVLKNASESLNSRMDRVEKKN
pIFNFH10
pIFNFH42
pIFNFH23
                  EDRLFESI
pIFNFH27
                  GDSLFDNIQSEEAN
pIFNFH14
                  KDRLFENTVRREKRI
pIFNFH01
                  EDRLFENTQRRQKKRNKKK
pIFNFH10
                  KYRLFENTQSEETKNNKKQ
pIFNFH11
                  EDRLFENTQRSQKKKNKKQ
pIFNFH13
                  KTAYLKIHRGDKRKKYKRMKHT
pIFNFH31
                  EDSLFENIQSEKKAKKVKQTNKKRSMY
pIFNFH03
                  RGGWITSSGVQDQPGQGSETSSLLKIQKLAGCSGRHL
DIFNEH15
                  DRLFENTKSEETNGKRIKCNEAHLQELENSFKMGNLKVIGLK
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WO 2004/050702 PCT/EP2003/050939

22/23

ABG00143	MGLRCDSETSWLQV	rfsti pavva	igtoffcprk	VEEKVK RTL	SCTSGTSSAT	esikwal					
ABG00143	GAPDSRTWLLDGISGPALGQRGAHCPQRHRQTSTSIKTIQEN										
pIFNFHcon INSP037 AAM70428 ABG00143	10 MTSPNELNKAPGT MTSPNELNKLPWTN MTSSNKLNKAPGTN	PGETEICDLS	DTEFKISVLK GEFKIAVLR	NLKEIQDNKE KLKEIQDNKE	KESRILSDKY KDFRILSDKF	KKQIEI					
pifnfhcon INSP037 AAM70428	70 IKKNQAEILELKNI IKGNQAEILELRNA IKKNQSEIQGLKNA	DGTL	FNSRIDQAEE	II							

Figure 23

